RAW SEQUENCE LISTING PATENT APPLICATION US/09/025,635

DATE: 03/06/98 TIME: 11:32:07

INPUT SET: S2851.raw

This Raw Listing contains the General Information Section and up to the first 5 pages

Sul Jal 1 SEQUENCE LISTING 2 General Information: (1) 3 4 (i) APPLICANT: Pang, Sheng-Zhi 5 Gonsalves, Dennis 6 7 Jan, Fuh-Jyh 8 (ii) TITLE OF INVENTION: DNA CONSTRUCT TO CONFER MULTIPLE TRAITS 9 ON PLANTS 10 11 (iii) NUMBER OF SEQUENCES: 18 12 13 14 (iv) CORRESPONDENCE ADDRESS: 15 (A) ADDRESSEE: Nixon, Hargrave, Devans & Doyle LLP (B) STREET: Clinton Square, P.O. Box 1051 16 JUN (C) CITY: Rochester 17 18 (D) STATE: New York (E) COUNTRY: U.S.A. 19 20 (F) ZIP: 14603 21 (v) COMPUTER READABLE FORM: 22 (A) MEDIUM TYPE: Floppy disk 23 24 (B) COMPUTER: IBM PC compatible 25 (C) OPERATING SYSTEM: PC-DOS/MS-DOS 26 (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 27 (vi) CURRENT APPLICATION DATA: 28 (A) APPLICATION NUMBER: 29 (B) FILING DATE: 30 31 (C) CLASSIFICATION: 32 (vii) PRIOR APPLICATION DATA: 33 34 (A) APPLICATION NUMBER: US 60/035,350 35 (B) FILING DATE: 19-FEB-1997 36 37 (vii) PRIOR APPLICATION DATA: (A) APPLICATION NUMBER: US 60/062,870 38 (B) FILING DATE: 21-OCT-1997 39 40 (viii) ATTORNEY/AGENT INFORMATION: 41 42 (A) NAME: Goldman, Michael L. (B) REGISTRATION NUMBER: 30,727 43 (C) REFERENCE/DOCKET NUMBER: 19603/1552 44 45

(ix) TELECOMMUNICATION INFORMATION:

46

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47 48 49	(A) TELEPHONE: (716) 263-1304 (B) TELEFAX: (716) 263-1600	
50 51	(2) INFORMATION FOR SEQ ID NO:1:	
52	(+) anathron attractmentanta	
53	(i) SEQUENCE CHARACTERISTICS:	
54	(A) LENGTH: 46 base pairs	
55 56	(B) TYPE: nucleic acid (C) STRANDEDNESS: single	
50 57	(C) STRANDEDNESS: STRIGTE (D) TOPOLOGY: linear	
58	(D) TOPOLOGI. IIIIeal	
59	(ii) MOLECULE TYPE: cDNA	
60	(II) HOLDCOLL IIII. COMI	
61		
62		
63		
64	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
65		
66	AGCTAATCTA GAACCATGGA TGACTCACTA AGGAAAGCAT TGTTGC	46
67		
68	(2) INFORMATION FOR SEQ ID NO:2:	
69		
70	(i) SEQUENCE CHARACTERISTICS:	
71	(A) LENGTH: 38 base pairs	
72	(B) TYPE: nucleic acid	
73	(C) STRANDEDNESS: single (D) TOPOLOGY: linear	
74 75	(D) TOPOLOGI: Timear	
76	(ii) MOLECULE TYPE: cDNA	
77	(II) MODDCODE IIID. CDIA	
78		
79		
80		
81	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
82		
83	TACAGTTCTA GAACCATGGT CTGGAAAACC TTGACCAG	38
84	(2) INFORMATION FOR CEO ID NO. 2.	
85 86	(2) INFORMATION FOR SEQ ID NO:3:	
87	(i) SEQUENCE CHARACTERISTICS:	
88	(A) LENGTH: 41 base pairs	
89	(B) TYPE: nucleic acid	
90	(C) STRANDEDNESS: single	
91	(D) TOPOLOGY: linear	
92		
93	(ii) MOLECULE TYPE: cDNA	
94		
95		
96		
97	(wi) GEOTHNOR DECERTION, GEO TO NO 3	
98 99	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	

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100 101	TACAGTTCTA GAACCATGGT AAAGCGATTT TACTTTTGGT A	41
102	(2) INFORMATION FOR SEQ ID NO:4:	
103		
104	(i) SEQUENCE CHARACTERISTICS:	
105	(A) LENGTH: 49 base pairs	
106	(B) TYPE: nucleic acid	
107	(C) STRANDEDNESS: single	
108	(D) TOPOLOGY: linear	
109		
110	(ii) MOLECULE TYPE: cDNA	
111		
112		
113		
114		
115	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:	
116	·	
117	AGATTCTCTA GACCATGGTG ACTTGATGAG CAAAGTCTGT GAGGCTTGC	.49
118		
119	(2) INFORMATION FOR SEQ ID NO:5:	
120		
121	(i) SEQUENCE CHARACTERISTICS:	
122	(A) LENGTH: 38 base pairs	
123	(B) TYPE: nucleic acid	
124	(C) STRANDEDNESS: single	
125	(D) TOPOLOGY: linear	
126		
127	(ii) MOLECULE TYPE: cDNA	
128		
129		
130		
131		
132	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:	
133		
134	TACAGTTCTA GAACCATGGA AAATACAAGG ATCTCGGG	38
135		
136	(2) INFORMATION FOR SEQ ID NO:6:	
137		
138	(i) SEQUENCE CHARACTERISTICS:	
139	(A) LENGTH: 41 base pairs	
140	(B) TYPE: nucleic acid	
141	(C) STRANDEDNESS: single	
142	(D) TOPOLOGY: linear	
143		
144	(ii) MOLECULE TYPE: cDNA	
145		
146		
147		
148		
149	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:	
150		
151	TACAGTTCTA GAACCATGGT AGAAGGGGAA AGAGTATGCT G	41
.152		

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			INPUI SEI: 52851.raw
153	(2)	INFORMATION FOR SEQ ID NO:7:	
154			
155		(i) SEQUENCE CHARACTERISTICS:	
156		(A) LENGTH: 36 base pairs	
157		(B) TYPE: nucleic acid	
158		(C) STRANDEDNESS: single	
159		(D) TOPOLOGY: linear	
160		(2)	
161		(ii) MOLECULE TYPE: cDNA	
162		(II) HOLLOOD IIID. ODMI	
163			
164			
165		/ //	
166		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:	
167			
168	AGC	ATTGGAT CCATGGTTAA CACACTAAGC AAGCAC	36
169			
170	(2)	INFORMATION FOR SEQ ID NO:8:	
171			
172		(i) SEQUENCE CHARACTERISTICS:	
173		(A) LENGTH: 36 base pairs	
174		(B) TYPE: nucleic acid	
175		(C) STRANDEDNESS: single	
176		(D) TOPOLOGY: linear	
		(D) TOPOLOGY: Tillear	
177			
178		(ii) MOLECULE TYPE: cDNA	
179			
180			
181			
182			
183		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:	
184			
185	TCT'	RGAGGAT CCATGGCTGA TCTTCATTCA TTTCAA	36
186			
187	(2)	INFORMATION FOR SEQ ID NO:9:	
188	(2)	1111 O101 11 1 1 1 1 1 1 1 1 1 1 1 1 1 1	
189		(i) SEQUENCE CHARACTERISTICS:	
190		(A) LENGTH: 36 base pairs	
191		(B) TYPE: nucleic acid	
192		(C) STRANDEDNESS: single	
193		(D) TOPOLOGY: linear	
194			
195		(ii) MOLECULE TYPE: cDNA	
196			
197			
198			
199			
200		(xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:	
201		(,	
202	արուր	TGAGGAT CCATGGATCC TGATATATAG CCAAGA	36
202	101	TOROGAL CCATOGATCC TORIATATAG CCAAGA	30
203	(2)	THEODMATION FOR CEO ID NO.10.	
	(2)	INFORMATION FOR SEQ ID NO:10:	
205			

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		INPUT SET: S2851.raw
206 207	(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 39 base pairs	
208	(B) TYPE: nucleic acid	
209	(C) STRANDEDNESS: single	
210	(D) TOPOLOGY: linear	
211		
212	(ii) MOLECULE TYPE: cDNA	
213		
214		
215		
216		
217	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:	
218	() 52232 2230 522 42 1.0 (20)	
219	TACAGTGGAT CCATGGTTAA GGTAATCCAT AGGCTTGAC	39
220		5,5
221	(2) INFORMATION FOR SEQ ID NO:11:	
222	(2) INFORMATION FOR BEG ID NO. 11.	
223	(i) SEQUENCE CHARACTERISTICS:	
224	(A) LENGTH: 36 base pairs	
225	(B) TYPE: nucleic acid	
225	(C) STRANDEDNESS: single	
226	(C) STRANDEDNESS: SINGLE (D) TOPOLOGY: linear	
	(D) TOPOLOGY: IIIIear	
228	(ii) MOLECULE EURE -DM	
229	(ii) MOLECULE TYPE: cDNA	
230		
231		
232		
233		
234	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:	
235		
236	TCTTGAGGAT CCATGGCTTA ATAACCTTCA TTATGC	36
237		
238	(2) INFORMATION FOR SEQ ID NO:12:	
239		
240	(i) SEQUENCE CHARACTERISTICS:	
241	(A) LENGTH: 36 base pairs	
242	(B) TYPE: nucleic acid	
243	(C) STRANDEDNESS: single	
244	(D) TOPOLOGY: linear	
245		
246	(ii) MOLECULE TYPE: cDNA	
247		
248		
249		
250		
251	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:	
252		
253	TCTTGAGGAT CCATGGAAAA GTCTTGAAGT TGAATG	36
254		
255	(2) INFORMATION FOR SEQ ID NO:13:	
256		
257	(i) SEQUENCE CHARACTERISTICS:	
258	(A) LENGTH: 50 base pairs	
	-	

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/09/025,635

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